

Amendments to the Specification

Please replace the paragraph at page 6, lines 3 through 9 with the following amended paragraph:

FIGS. 1A and 1B are schematic representations of the predicted secondary structure of the miR15 (SEQ ID NO:1) and miR16 (SEQ ID NO:2) precursor RNA, respectively. The RNA secondary structure prediction was performed using the “mfold” program, version 3.1 of Matthews et al. (1999), *J. Mol. Biol.* 288:911-940, and manually refined to accommodate G/U wobble base pairs in the helical segments. The sequence of the processed miR15 (SEQ ID NO:3) and miR16 (SEQ ID NO:4) miRNA is underlined. Adapted from Lagos-Quintana et al. (2001), *Science* 294:853-858.